

## SEQUENCE LISTING

&lt;110&gt; Garber, Mitchell Ehren

<120> METHODS AND COMPOSITIONS FOR USE IN  
EVALUATING AND TREATING NEOPLASTIC DISEASE CONDITIONS

&lt;130&gt; STAN-349WO

&lt;150&gt; 60/629,527

&lt;151&gt; 2004-11-18

&lt;150&gt; 60/558,953

&lt;151&gt; 2004-04-02

&lt;160&gt; 13

&lt;170&gt; FastSEQ for Windows Version 4.0

&lt;210&gt; 1

&lt;211&gt; 20

&lt;212&gt; PRT

&lt;213&gt; human

&lt;400&gt; 1

Met	Glu	Ala	Ala	Asp	Ala	Ser	Arg	Ser	Asn	Gly	Ser	Ser	Pro	Glu	Ala
1				5					10					15	
Arg	Asp	Ala	Arg												
			20												

&lt;210&gt; 2

&lt;211&gt; 17

&lt;212&gt; PRT

&lt;213&gt; human

&lt;400&gt; 2

Glu	Leu	His	Leu	Lys	Pro	His	Leu	Glu	Gly	Ala	Ala	Phe	Arg	Asp	His
1				5					10					15	
Gln															

&lt;210&gt; 3

&lt;211&gt; 18

&lt;212&gt; PRT

&lt;213&gt; human

&lt;400&gt; 3

Glu	Gly	Glu	Gly	Leu	Gly	Gln	Ser	Leu	Gly	Asn	Phe	Lys	Asp	Asp	Leu
1				5					10					15	
Leu	Asn														

&lt;210&gt; 4

&lt;211&gt; 17

&lt;212&gt; PRT

&lt;213&gt; human

&lt;400&gt; 4

Arg Glu Thr Ile Pro Ala Lys Leu Val Gln Ser Thr Leu Ser Asp Leu  
1 5 10 15  
Arg

<210> 5  
<211> 17  
<212> PRT  
<213> human

<400> 5  
Asp Pro Ala Lys Val Gln Ser Leu Val Asp Thr Ile Arg Glu Asp Pro  
1 5 10 15  
Asp

<210> 6  
<211> 15  
<212> PRT  
<213> human

<400> 6  
Arg Val Ala Ala Lys Arg Leu Lys Glu Gly Asp Thr Met Met Gly  
1 5 10 15

<210> 7  
<211> 19  
<212> PRT  
<213> human

<400> 7  
Lys Thr Val Glu Ser Leu Glu Glu Thr Leu Lys Lys Ala Ser Pro Asp  
1 5 10 15  
Gly Tyr Asp

<210> 8  
<211> 17  
<212> PRT  
<213> human

<400> 8  
Thr Thr His Ser Ile Ser Asp Gly Lys Asp Leu Glu Lys Leu Leu Thr  
1 5 10 15  
Glu

<210> 9  
<211> 24  
<212> PRT  
<213> human

<400> 9  
Glu Tyr His Lys Val His Gln Met Met Arg Glu Gln Ser Ile Leu Ser  
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Pro Ser Pro Tyr Glu Gly Tyr Arg  
20

<210> 10  
<211> 23  
<212> PRT  
<213> human

<400> 10  
Arg His Gln Leu Leu Cys Phe Lys Glu Asp Cys Gln Ala Val Phe Gln  
1 5 10 15  
Asp Leu Glu Gly Val Glu Lys  
20

<210> 11  
<211> 19  
<212> PRT  
<213> human

<400> 11  
Asp Leu Glu Val Lys Asp Trp Met Gln Lys Lys Arg Arg Gly Leu Arg  
1 5 10 15  
Asn Ser Arg

<210> 12  
<211> 30  
<212> DNA  
<213> human

<400> 12  
ggatccatgg aagctgcaga tgcctccagg 30

<210> 13  
<211> 71  
<212> DNA  
<213> human

<400> 13  
accggtgtgc atcctccgcc gccgcatacct ccgccgccgg cggctggggc ttcgttggac 60  
ccaatcccgt t 71